

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/537,839
Source: PCT/10
Date Processed by STIC: 6/16/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/24/05

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Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/537,839 //

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 Wrapped Nucleic
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 Misaligned Amino Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 Variable Length Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 PatentIn 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 Skipped Sequences (OLD RULES) Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 Skipped Sequences (NEW RULES) Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 Invalid <213> Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 Use of <220> Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12 PatentIn 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,839

DATE: 06/16/2005
TIME: 10:18:53

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Output Set: N:\CRF4\06162005\J537839.raw

3 <110> APPLICANT: THE CORPORATION OF THE TRUSTEES OF THE ORDER OF THE SISTERS OF MERCY

IN

4 QUEENSLAND
6 <120> TITLE OF INVENTION: NOVEL THERAPEUTIC MOLECULES AND USES THEREOF
8 <130> FILE REFERENCE: 12381870/TDO
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/537,839
C--> 10 <141> CURRENT FILING DATE: 2005-06-06
10 <150> PRIOR APPLICATION NUMBER: 2002953223
11 <151> PRIOR FILING DATE: 2002-06-12
13 <160> NUMBER OF SEQ ID NOS: 32
15 <170> SOFTWARE: PatentIn version 3.1

pp 12-13

*Does Not Comply
with Diskette Needs*

ERRORED SEQUENCES

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4242	taataccacc	acccaaatcca	ccctcacacc	ttaccaaatac	tttactacca	aatttttattt	26280
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4246	aatatttgaa	aagtatcaat	ggctctccct	cacttaaca	aataatttgg	ccaaattttct	26400
4248	ctcactaatac	ttaaattctt	ccaggtctca	tttaacaag	caccagttga	aataatgggt	26460
4250	attatgaata	tattaataa	taaaaaatct	gtcaaaataa	ttctgaaaag	tttcacatt	26520
4252	ctttagagtc	gattactata	cataatacaa	gtacctgccc	aaaagtatgt	ttgttctcc	26580
4254	aaaccaggtt	atttcctca	atttattaaa	ctggattctt	atagaaagat	tacaagggtt	26640
4256	tgaataggca	agaactgca	agttgtgcat	acttacagaa	atttagggaa	acacacaaaag	26700
4258	aatgaatgaa	tgcaaagaca	cacaaaacca	ggtactttac	tgccaggtctt	gggtttttt	26760
4260	tcagttaaac	aaatcttaaa	tcccaaagc	caggtgaacc	attaaacctc	aataatgtaa	26820
4262	cacattatct	aaatactaga	attcctaaag	ggaacagtag	gtctgaattt	gtagtcttag	26880
4264	atcagaaagt	aaagcgaagg	aagacaaact	ggagcatggt	atatacttc	gtgatttcca	26940
4266	ctatgggtc	ttcagtagag	tggagaagt	cacaacttcc	tagaaaattt	gagtactatt	27000
4268	cttcaggct	agaaatataa	atcttggtag	taacaaatgc	gaaattttt	gtttctcat	27060
4270	cagtgtggg	aattctcaac	acagagaaaat	gttacacaa	aactacagtt	gtgctcatca	27120
4272	aaatataactt	gcaacttgt	taccctaatt	tcccaaatac	tctttgtct	tttcctttt	27180
4274	agataaccac	attttaatat	cagcatttgt	gattgcttagc	acggttaattt	tgacagttt	27240
4276	gggagcaatc	atttggttcc	tgtacaaaaaa	acattctgat	tctcggttca	ccacagttt	27300
4278	ttcaaccgca	ccccaaatcac	cttataatga	agactgtgtt	ttggtagtt	gagaagaaaa	27360
4280	tgaatatct	gttcaattt	actaagttt	ttgttaatctt	gcactaagac	atcaacaaa	27420
4282	tgcctggca	gagataactt	gggaaagatt	ttaatataaa	acttgacatt	ggatattaga	27480
4284	gtttatgg	tattcctt	tccagtaaca	tttttatgt	ctcatctgt	gtaaaaagtc	27540
4286	tttaggttca	ttaaaaaaaac	aggttttaga	aatgatctt	gatctaata	atgtatttt	27600
4288	agcatccgt	caaaggcaga	atctgtact	tgaatgaagg	aaagctaaa	gcccaagcag	27660
4290	ataaaaataa	aagcccagcc	tatttgc	gcctgctgt	tctccctat	ttagttgacc	27720
4292	cacttagtt	tatatgtt	ttagtaaca	tgaatgggg	aataagtgt	ttaagtaca	27780
4294	tcccatacat	ttaaatatct	ttgataattt	ttatttttt	ggcagataat	tcctctagaa	27840
4296	tgtgtatctt	tttatgattt	agatgaagaa	aattttacaa	cttttaacac	cccacaccaa	27900

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Input Set : A:\pto.da.txt
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4298	tttagttc	attacttta	cacacaccat	tttatcacaa	atgactcaag	tttaatgaa	27960
4300	tgttataaa	ttatggaaa	caaaatatga	tcgctgtgtc	caggatggca	tagagaaagc	28020
4302	tggcaattag	gttaacactt	acatattata	gtgcccctt	aaggatttct	ctttgccac	28080
4304	cataccttt	gtactttccc	ctataacaaga	tgtatctcat	tctcctcaag	catttataaa	28140
4306	ttttccttc	aatgacatga	aaactgtgca	agcaaaaacc	gaagaaaaac	acttaagtac	28200
4308	aactgttagt	acagtgtca	aagtttcag	tgcatattt	gtacatttta	agaaaaaagg	28260
4310	gaaaatcatt	tggggagtaa	aaaaatgaaa	aagctgaaac	gagtaattt	cctcaccatc	28320
4312	aataaaccaa	aaacaggaaa	gataaagaat	gtataaattt	cacgtaaattt	agtcacgtat	28380
4314	cacttatcaa	tggggatacg	ttctaagaaa	tgcatagtt	ggaatctt	tgtgaaaatc	28440
4316	agcttgtatt	tacacaaacc	cagatggtag	agcttattt	gtcccaaacc	tacacagcat	28500
4318	gttactgtgc	tgaataactgc	agacaattgt	aacacaatat	ttgtgtatct	aaatataaaaa	28560
4320	aaggtagact	aaaaatatgg	tctactaagg	aaacactgtt	ctatatgtgg	tcattactg	28620
4322	actgaagtagt	actgtctaga	agtctgaggc	tcaaagaaaa	gtaatccctc	ttctgaatcc	28680
4324	acaccccatc	aattatctta	cttcttctg	gggagataga	tagatatact	atctcaactag	28740
4326	cttgactaat	ggcaacaaag	ttccagctt	tgttagtctt	ttttatttgc	cacatgaatc	28800
4328	gaaaacactc	atcacaatta	atggcactat	cattaatgag	acatgagtaa	ctaaaaaagg	28860
4330	atagaaaact	attaacagtg	cggctacatg	gtactgaaaa	tgcagggatt	acaccagctg	28920
4332	ttacacaagc	acaagcatgc	tctgttaagag	ctttacattt	ctgagattt	gtatagtgtat	28980
4334	tgagatgtct	attttattat	tgatagacta	ttactaatgt	caatattgaa	cactaccctg	29040
4336	gaattcctgc	ctggtttcc	tacccaaatt	gtaccactcc	ttgaagaact	acaggcacag	29100
4338	taaaaaaaat	atggcgtatt	atgtgaacta	aaagagttt	aaaggagttc	ttaaaggagg	29160
4340	ggtagaattt	gggttagaaaa	gtgattaagt	ccaaacttaaa	accaacagtc	tcaaacgtot	29220
4342	acaactacaa	tgtccaaatga	gccactagcc	acatgaggct	atthaagtaa	atttagttt	29280
4344	aaatccagtt	ttcgaattac	attagccaca	ttgtcaagtg	ttcaaatcac	aggtggtag	29340
4346	tggctactgt	actgggcaac	atacattata	gaacatttc	attataggaa	gttttatttg	29400
4348	gcagtgtgc	tcttaatcc	taccttccac	tcaactccca	tacaacttcc	ttttgtacat	29460
4350	tttgataactt	tctacctaatt	ggcagcttt	ccaaaatagc	tgctttaaac	tctgatttaa	29520
4352	tttcaatata	ttggtttcat	tttcaacag	gccaagaggc	ctctggtaat	gaagtgtat	29580
4354	atatatataat	atgatggagt	ctcaactgtgc	tgcccaggct	acagtgcagt	ggctcgatct	29640
4356	tggctctctc	caatctccgc	cttgcagggtt	ttcaagcaat	tctcctgcct	cagcctcctt	29700
4358	agtagctggg	accacagaca	tctgtcacca	cacccagcta	actttttgt	tttttggtag	29760
4360	agacggggtt	tcgccccattt	gactgggctg	gtctcaaact	cctgacctca	agtgtatccac	29820
4362	ccaccttgg	ctcccaaagt	gctgggatta	catgcgtgag	ccaccacact	tggcctacat	29880
4364	ttttcttta	tataccagaa	catctataac	aggcacctt	tctactcatt	agtgaagaga	29940
4366	taattggatt	acacaggcg	gcttggttac	tacatccaga	atgtagaaac	tgctttcttc	30000
4368	aacatcttgg	ttctagctag	taataacaat	ataattctt	ggcagatatt	cagaataaca	30060
4370	ttttaaacta	cattttctta	gaaaattgca	ttctttagt	gagcagtgt	tggctcttt	30120
4372	tgttcagaat	ttaaaaactga	taaccaatga	aagcctttc	tcttatttct	ctaccgtcat	30180
4374	ttacatgata	atctgaagct	aatatgacaa	tattttaaata	ctaagtggta	ctagggaaact	30240
4376	acaagaatac	tgtaaagctt	aagccattgt	tatcaactgtc	attagcatt	taataacaaa	30300
4378	actatacaga	attatgtgca	taccaatgaa	tgtttgtac	catctagtt	atttttttaa	30360
4380	ataaaagtttt	atgggttaag	cagaagacaa	ctgtcatact	gaatttttatt	aaaagtataat	30420
4382	atacttcaaa	ttcaaagcat	cccttaggac	ccacacagaata	tattttttact	accaccctt	30480
4384	aattttatata	ttttcttta	agacagacaa	tgcaaaggta	actggcaaga	ggtgagcaaa	30540
4386	tgttttagaa	catttatatt	attgcttaaa	atgagattt	aaattgtat	aaaattctt	30600
4388	gttatgaagt	ctgatgtctt	ctttgagcac	cagttttaaa	ggaaacattt	caaacagtaat	30660
4390	aataaaatcag	tgtgtacttt	attttgagcc	aatgttttac	ataagaccaa	caaactaagt	30720
4392	gtgggcaaac	tgttcaatca	cttttgagaa	attacatgg	tccttatttag	gtttgaaaaaa	30780
4394	gtagtttgg	gccataacat	cagaccttaa	ggtttttaa	agtgtatctt	aagggtactg	30840

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Input Set : A:\pto.da.txt
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4396	acagaagaaa aatacagtgt tatggaagta tacacaactc atgaccacac agtata	30900
4398	cattcatggt gaataatgtt catctgttct ccaaaggtag taaaaatata taagtcatca	30960
4400	aataaaatgtc cagaattacc aaggaacagt taaaaaggac caggatccaa acaggattt	31020
4402	tatataattta tctataaaaca agtgacaaca ctaaaaaaca cattgaaaac cagtgttt	31080
4404	aactctgaat gtgggattta aaaatattcc attaacttga gattgcaata attttttt	31140
4406	accaaatacg atttgactgg caagtctcat ataatttaca gctgtatata tcctaaaagg	31200
4408	aatatctaca tactatagcc tatcaaatat aatctgcatt ttgcaaatca ataca	31260
4410	ctatccatca tttatatacatt ttatatttat atactttta aaaatttagat ttatttgcta	31320
4412	ggtaatttg gaattcaaat taaatttgga aacttaataa tgctcttcata accttct	31380
4414	ccccattttt ttttaattt ttaaacactg gaagcaatg attatcaaaa tgagactgca	31440
4416	acttaaggca ttttaaaaaga aaaaataatt aggtgcata tagcatttt ttcaaaagt	31500
4418	atattttgtc ccacttttct cactagcaag agtaaaacac aaacctttt tcataaata	31560
4420	gactacagta atcataaacac aaaaagggt tgataggcat tgcttagata tttaaaaca	31620
4422	ggtaataact ttcccactca cctaaaagaa aaaacctttt tgattaccag ttataaaaca	31680
4424	tcggatttgc tatgtaaaaa agtccagcag aattttaatt cagcaacact ggagaatgaa	31740
4426	tatataatata caggcataca tttgtatgtatatttatttttataatag tcaatgtatt	31800
4428	tattaggcat tcccccacaa aagtaatttattttatgttccatattttaa taaacactta	31860
4430	tgttcacaga tcatccatct gtcttatatg aagttaggca atatcaatg tcctgtt	31920
4432	gtctccgtct tcttcggaat catcctctga agtgttggaa agaaaacagg atgatcaaga	31980
4434	atgattttaa gggggataag aatgggaagt gtaaatatga agccagaaca gcaatgactt	32040
4436	gtaaaaacaa aaatcccaat tttagtatattt gttgtatcaa aatgataagt tttatgacag	32100
4438	ggtaggggt gtctccgtgt ggggtatacc caaccacaat tctcccgcc cccacaccgt	32160
4440	tttgtatgt tatacaatttttataaaaaa tggaaagtgtct taaaaatgtg gttcaagagc	32220
4442	atcttctaaa atgtttctga gatcaatctg ttttcatagt agctaatgtat atataa	32280
4444	tagaaaaaaa gtcaacttaa ggcaagtgtct ctcacacacc tgagtaaaaat ctggctt	32340
4446	accgctgcat gttttgtca tccgatgaga atgaaaagaa gtgtacatag aataagaggt	32400
4448	tatgggtgtt cttgcaattt acatcaaagc tatgtctata caatgccaag ataaaagt	32460

E--> 4457 - 3 -

delete

see p. 13 for more errors

10/537,839 13

<210> 23
<211> 21
<212> DNA
<213> synthetic

<400> 23
gaccatggag cgAACATGAT a

invalid <213> response
see item 10 on
Error Summary
sheet

21

The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

VERIFICATION SUMMARY
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Input Set : A:\pto.da.txt
Output Set: N:\CRF4\06162005\J537839.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:28 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:25
L:1231 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:1229
L:1609 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:1606
L:1810 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:1807
L:1812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:48
M:341 Repeated in SeqNo=10
L:1901 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:2010 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:2040 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:2037
L:2069 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:2067
L:2096 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:18,Line#:2094
L:2125 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:20,Line#:2123
L:4457 M:254 E: No. of Bases conflict, this line has no nucleotides.

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